NUCLEOTIDE AND AA SEQUENCE

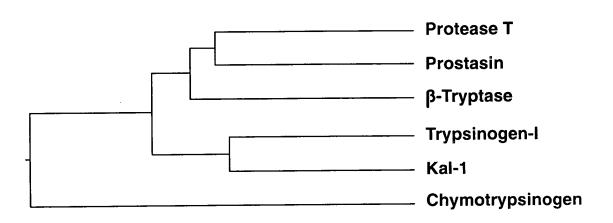
Nucleic acid sequence of Protease T (SEQ.ID.NO.:1)

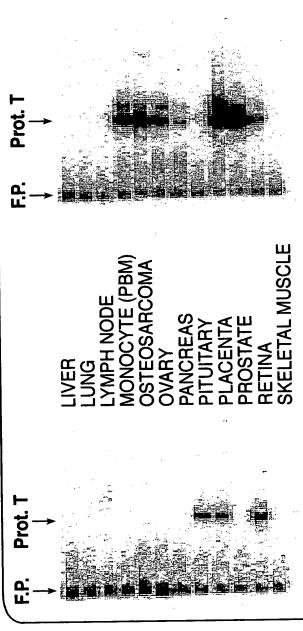
TCCTGCTGCTGTGTTTTTGGGTCTCAGAGGGCCAAGGCAGCAACAGCCTGTGGTCGCC CCAGGATGCTGAACCGAATGGTGGGCGGGCAGGACACGCAGGAGGGGCGAGTGGCCCTGGC AAGTCAGCATCCAGCGCAACGGAAGCCACTTCTGCGGGGGCAGCCTCATCGCGGAGCAGT GGGTCCTGACGGCTGCGCACTGCTTCCGCAACACCTCTGAGACGTCCCTGTACCAGGTCC AGGTGGAGAGCAACCCCCTGTACCAGGGCACGGCCTCCAGCGCTGACGTGGCCCTGGTGG AAGACCTCCTGCCCGAACCGCGGATCCTGCAGAAACTCGCTGTGCCCATCATCGACACAC CCAAGTGCAACCTGCTCTACAGCAAAGACACCGAGTTTGGCTACCAACCCAAAACCATCA AGAATGACATGCTGTGCGCCGGCTTCGAGGAGGGCCAAGAAGGATGCCTGCAAGGGCGACT CGGGCGCCCCTGGTGTGCCTCGTGGGTCAGTCGTGGCTGCAGGCGGGGGGTGATCAGCT GGGGTGAGGGCTGTGCCCGCCAGAACCGCCCAGGTGTCTACATCCGTGTCACCGCCCACC AGAAGTGAGACCCCCGGGGCCAGGAGCCCCTTGAGCAGAGCTCTGCACCCAGCCTGCCCG CCCACACCATCCTGCTGGTCCTCCCAGCGCTGTTGCACCTGTGAGCCCCACCAGACT CATTIGTAAATAGCGCTCCTTCCTCCCCTCTCAAATACCCTTATTITATGTTTCTC CCAATAAAAACCCAGCCTGTGTGCCAGCTG

>Amino Acid Sequence of Protease T (SEQ.ID.NO.:7)

MRRPAAVPLILLLCFGSQRAKAATACGRPRMLNRMVGGQDTQEGEWPWQVSIQRNGSHFC GGSLIAEQWVLTAAHCFRNTSETSLYQVLLGARQLVQPGPHAMYARVRQVESNPLYQGTA SSADVALVELEAPVPFTNYILPVCLPDPSVIFETGMNCWVTGWGSPSEEDLLPEPRILQK LAVPIIDTPKCNLLYSKDTEFGYQPKTIKNDMLCAGFEEGKKDACKGDSGGPLVCLVGQS WLQAGVISWGEGCARQNRPGVYIRVTAHHNWIHRIIPKLQFQPARLGGQK

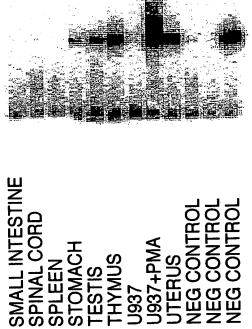
FIG. 2





SUBTHALMIC NUCLEUS BREAST CARCINOMA BONE MARROW AMYGDALA CAUDATE NUCLEUS CEREBELLUM CEREBRAL CORTEX SUBSTANTIA NIGRA **BCELL LYMPHOMA** HIPPOCAMPUS HYPOTHALAMUS **THALAMUS**

3/6



SUMYH.

1937

ESTIS

SPLEEN

TERUS

JURKAT+PMA+PHA-M KERATINOCYTE FETAL BRAIN FIBROBLAST ENDOTHELIA **EUKOCYTE EPIDERMIS** FAT CELL JURKAT KIDNEY HEART COLON

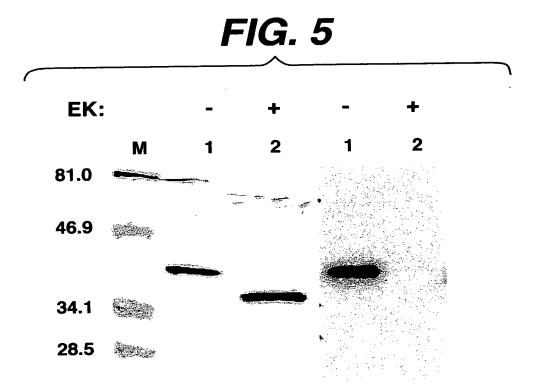
FUSION PROTEIN

Nucleic Acid sequence of PFEK-PROTT-HIS fusion protein (SEQ.ID.NO.:8).

GAATTCACCACCATGGACAGCAAAGGTTCGTCGCAGAAATCCCGCCTGCTCCTGCTG GTGGTGTCAAATCTACTCTTGTGCCAGGGTGTGGTCTCCGACTACAAGGACGACGACGAC GTGGACGCGCCCCTTTGCTGCCCCCTTTGATGATGATGACAAGATCGTTGGGGGCTAT GCTCTAGAGGAGGGCGAGTGGCCCTGGCAAGTCAGCATCCAGCGCAACGGAAGCCACTTC TGCGGGGCAGCCTCATCGCGGAGCAGTGGGTCCTGACGGCTGCGCACTGCTTCCGCAAC ACCTCTGAGACGTCCCTGTACCAGGTCCTGCTGGGGGCAAGGCAGCTAGTGCAGCCGGGA CCACACGCTATGTATGCCCGGGTGAGGCAGGTGGAGAGCAACCCCCTGTACCAGGGCACG GCCTCCAGCGCTGACGTGGCCCTGGTGGAGCTGGAGGCACCAGTGCCCTTCACCAATTAC ATCCTCCCCGTGTGCCTGCCTGACCCCTCGGTGATCTTTGAGACGGGCATGAACTGCTGG GTCACTGGCTGGGCCAGCCCCAGTGAGGAAGACCTCCTGCCCGAACCGCGGATCCTGCAG AAACTCGCTGTGCCCATCATCGACACACCCAAGTGCAACCTGCTCTACAGCAAAGACACC GAGTTTGGCTACCAACCCAAAACCATCAAGAATGACATGCTGTGCGCCGGCTTCGAGGAG GGCAAGAAGGATGCCTGCAAGGGCGACTCGGGCGCCCCCTGGTGTGCCTCGTGGGTCAG TCGTGGCTGCAGGCGGGGTGATCAGCTGGGGTGAGGGCTGTGCCCGCCAGAACCGCCCA GGTGTCTACATCCGTGTCACCGCCCACCACAACTGGATCCATCGGATCATCCCCAAACTG CAGTTCCAGCCAGCGAGGTTGGGCGGCCAGAAGTCTAGACATCACCATCACCATCACTAG CGGCCGCTTCCCTTTAGTGAGGGTTAATGCTTCGAGCAGACATGATAAGATACATTGATG AGTTTGGACAAACCACAACTAGAATGCAGTGAAAAAAATGCTTTATTTGTGAAAATTTGTG ATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTGAC

Amino Acid sequence of PFEK-PROTT-HIS fusion protein (SEQ.ID.NO.:9).

MDSKGSSQKSRLLLLLVVSNLLLCQGVVSDYKDDDDVDAAALAAPFDDDDKIVGGYALEE GEWPWQVSIQRNGSHFCGGSLIAEQWVLTAAHCFRNTSETSLYQVLLGARQLVQPGPHAM YARVRQVESNPLYQGTASSADVALVELEAPVPFTNYILPVCLPDPSVIFETGMNCWVIGW GSPSEEDLLPEPRILQKLAVPIIDTPKCNLLYSKDTEFGYQPKTIKNDMLCAGFEEGKKD ACKGDSGGPLVCLVGQSWLQAGVISWGEGCARQNRPGVYIRVTAHHNWIHRIIPKLQFQP ARLGGQKSRHHHHHH



Protease: PFEK-Protease T-HIS

Protease Activity of Protease T

H-D-Pro-HHT-Arg-pNA
H-D-Lys(CBO)-Pro-Arg-pNA
H-DL-Val-Leu-Arg-pNA

